1/30

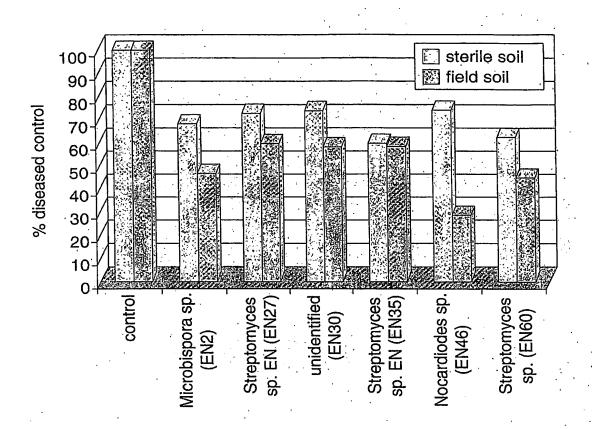
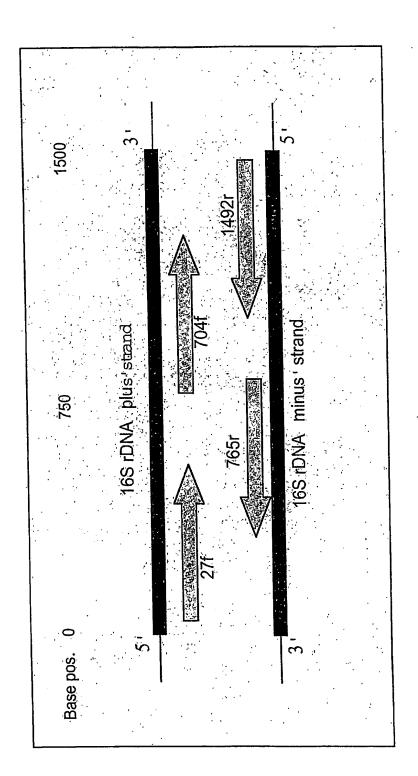


Figure 1





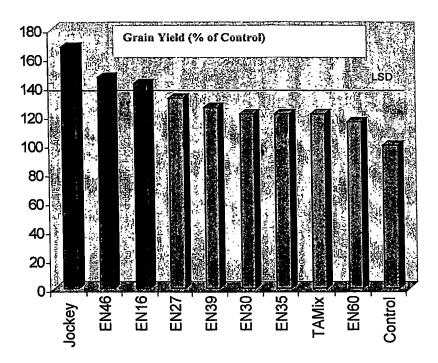
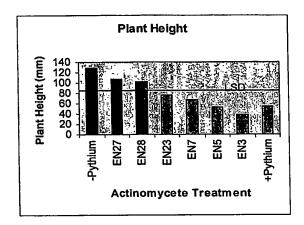


Figure 3

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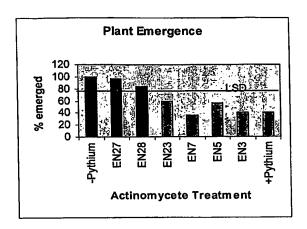


Figure 4

Figure 5

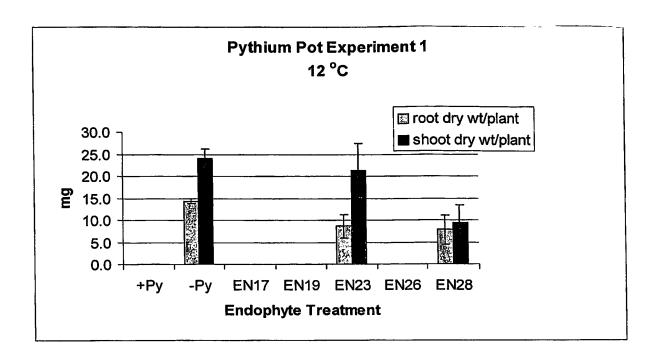


Figure 6

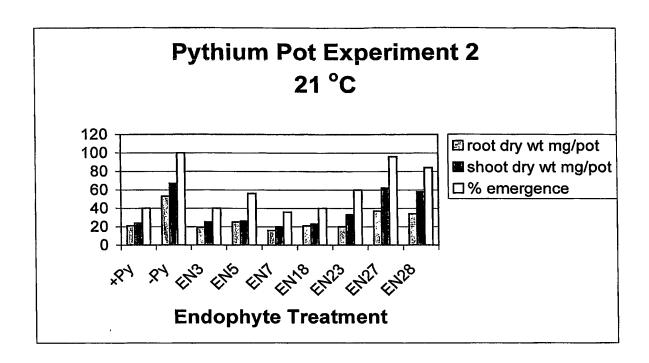
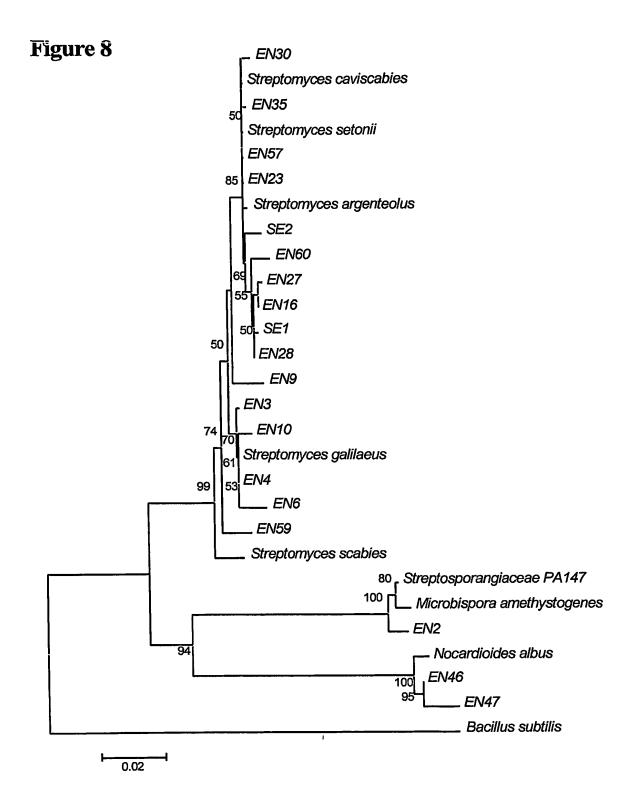


Figure 7



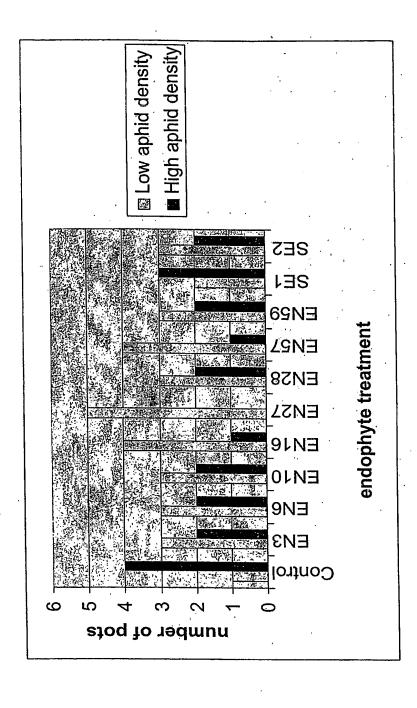


Figure 9

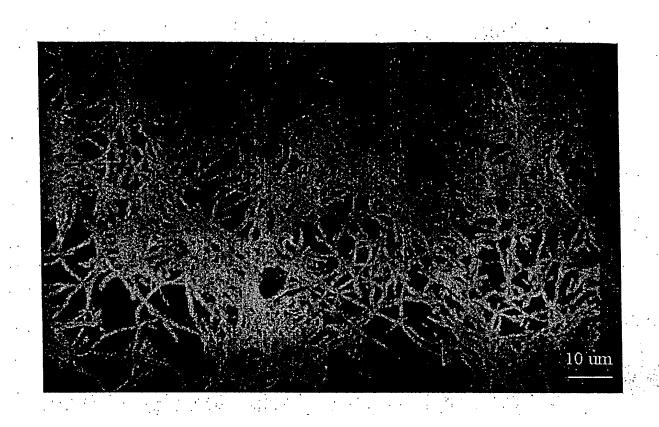
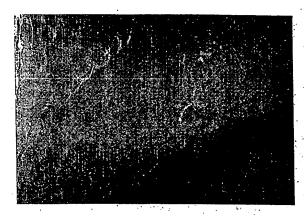


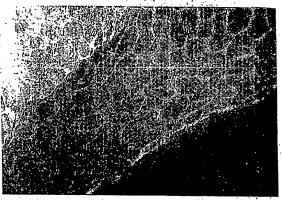
Figure 10

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11.1 Blue excitation/ green emission

11.2 UV excitation/ blue emission





11.3 Image enhanced merge

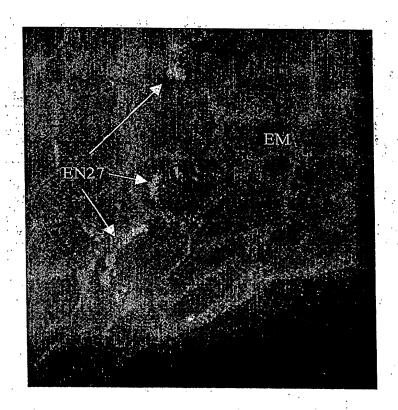
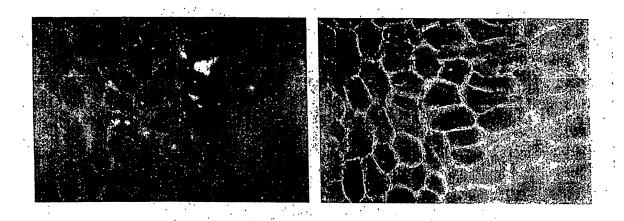


Figure 11

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12.1 Blue excitation/ green emission

12.2 UV excitation/ blue emission



12.3 Image enhanced merge

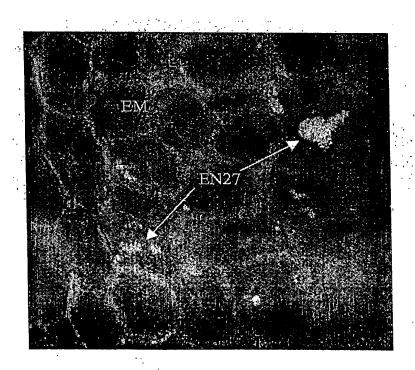


Figure 12

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13.1 Blue excitation/ green emission 13.2 UV excitation/ blue emission



13.3 Image enhanced merge

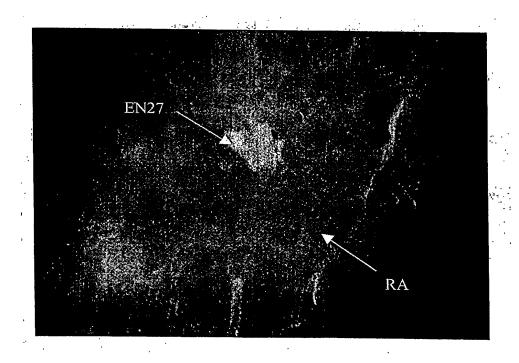


Figure 13

14.1 Blue excitation/ green emission 14.2 UV excitation/ blue emission



14.3 Image enhanced merge

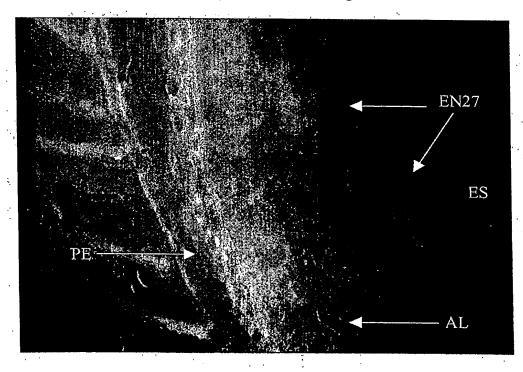


Figure 14

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FIGURE 15

EN2

SEQ ID NO:1

CTTAACACATGCAANTCAAGCGGAAAGGCCCTTCGGGGTACTCAANCGGCNAACGGGTGATTAACACNTGANTAA CCTGCCCTGACTCTGGGATAANCCTGGGAAACTGGGTCTAATACCGGATACAACCATTTCTCNCATGGGATGGT GGTGGAAANTTTTTNCGGTTGGGGATGGGCTCGCGGCCTATCACCTTGTTGGTGGGGTGATGGCCTACCAAGGCg ACGAACGGTagCCCGCCTGAgAGGGCGACCGGCCaCaCTGGGACTGAGACaCCGCCCGAACTCCTaCgGGAGGCA $\tt gCACTgGGGGAATaTTGCCCATGGGCGGAAGCCTGACGCAGNGACGCCGCGTGGGGGGATGACGGCCTTNGGGTTGT$ AAACCTNTTTCAGCAGGGACGAAGTTGACGTGTACCTGTAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGC GGTAATANGTAGGGCGCGAGCGTTNTCCGGAATTATTGGGCGTAAAGAGTTTGTAGGTGGCTTGTTGCCTTTGCC GTGAAAGCCCGTGGCTTAANTACGGGTTTGCGGTGGATACGGGCAGGCTAGAGGCTGGTAGGGGCAAGCGGAATT $\verb|CCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAGACACCGGTGGCGAAGGCGGCTTGCTGGGCCAgTTCtga| \\$ $\tt CGGTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCTGTAAACGTTGGGCGCTA$ GGTGTGGGGGGTCTTCCACGATCTCTGTGCCGTAGCTAACGCATTAAGCGCCCCGCCTGGGGAGTACGGCCGCAAG GCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCGGGGGGCGTTGCTTAATTCGACGCAACGCGAAGA ACCTTACCAAGGTTTGACATACACCGGAAACACTCANANATGGGTGCCTCCTTTGGACTGGTGTACAGGTGGTGC ATGGCTGTCNNCACCCTCGTGTCGTNAGATGTNGGGTTAAGTCCCGCAACGANCGCAACCCTTGGTTCCATGTTG CCAGCACNCCCTTTGNGGTGGTGGGGACNCATGGGGANAATGCCGGGGTCNACTCNGGAGGAAGGTGGGGATGACG TCAAGTNATCNTGCCCCTTATGTTCTTGNNGTG

EN3

SEQ ID NO:2

GCTGGCGGCGTGCTTAACACATGCAAGTCGAACGATGAACCACTTCGGTGGGGATTAGTGGCGAACGGGTGAGTA ACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTNCTGCTC TCATGGGCAGGGGTTAAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTTGGTGAGGTAATGGC TCACCAAGGCGACGGCTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCC TACGGGAGGCAGCAGTGGGGAATATTGCaaCAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACG ${\tt GCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAC}$ TACGTGCCAGCAGCCGCGTAATACGTAgGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTtGTAgGC GGCTTGTCACGTCGGGTGTGAAAGCCCGGGGCTTAACCCcGGGTCTGCATTCGATACGGGCTAqCTAqAGTGTGG TAGGGGAGATCGGAATTCcTGGTGTAGCGGTGAAATGCqCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGaT ctctgggccattactgacgctgaggagcgaaagcgtgggagcgaacaggattagataccctggtagtccacgc GTAAACGGTGGGAACTAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCATTAAGTTCCCCGCCT GGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCAGCGGAGCATGTGGCTTAA TTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACACCGGAAAGCATCAGAGATGGTGCCCCCCTTGTGG TtCGGTGTACAGGTGGTGCATGGCTGTCGTCAGCTCGTGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCA ACCCTTGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACCGCCGGGGTCAACTCGGAG GAAGGTGGGGACGTCAAGTCATCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGcCGGTACAAA GAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTcTGCAACTCGAcCC CATGAAGTCGGAGTTGcTAATAATCgCANATCAgCATTGCTGCGGTGAATACGTTcCCGGGCCTTGTACAcACCG CCcGTCACGTcACGAAAGTCGgTAAcACCcGAAgCCGGTGGCCAACCCCTTgTGGGAGGAGCTGTCGAAGGTGG GACTGGCGATTG

15/30

[FIGURE 15 CONTINUED]

EN16

SEQ ID NO:7

GCTTNTTGGTGGGNCNATGGCCTACCAAGGNGAGGACGGNTANCCNGCCTGNGAGGGAGACCGNCCACACTGGGA
ATGNGANACGGCCCAGAATCCTACGGGAGGCAGCANNGGGGAANATTGCACAANGGGCGAAAGCCTGATGCAGNG
ANGCCGCGTGAGGGAAGACGGCCTTTGGGTTGTAAACCTNTTTNAGCAGGGAAGAAGCGAAAGTGACGGTACCTG
CAGAAGAAGCGCCGGCTAANTANGTGCCAGCAGCCGCGGTAATANGTAGGGCGCAAGCGTTGTCCGGAATTATTG
GGCGTAAAGAGCTTGTAGGCGGCTTGTCANGTNGGATGTGAAAGCCCGGGGCTTAACCCCGGGTTTGCATTTGAT
ACGGGCTAGCTAGAGTGTGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAA
CACCGGTGGCGAAGGCGGATCTCTGGGGCCATTACTGACGCTGAGGAGCGAAAAGCGTGGGGAGCGAACAGGATTAG
ATACCCTGGTAGTCCACGCCGTAAACGTTGGGAACTAGGTGTTGGCGACATTCCACGTCGTCGTCGCCGCAGCTA
ACGCATTAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAG
CAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAG
AGATGGTGCCCCCCTTGTGGTCGGTATACAGGTGGTCATGCCTTCGGGGTGATGTTGGGTT
AAGTCCCGCAACGAGCCAACCCTTGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGAC
TGGGGGTCTGNAACTCGACCCCATGAANTCGGAGTTGCTAATAATCCCAAATTCANCATTGGTGCGGTGAATACT
TCCCGGGCCTGGTACACNACCGCCCGTCAACTCACGAAAGTCGGTNAAACCCGAAACCGGTGGGCCAACCCCTTG
TGGGAAGGAACTGGCCTAAACTGGCGATTGGGAC

EN23

SEQ ID NO:10

ACGAACGCTGGCGGCGTGCTTAACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACG GGTGAGTAACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACAC TCTGTCCCGCATGGGACGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTTGGTGGG GTAATGGCCTACCAAGGCGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCC CAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGG GATGACGGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCC GGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCT AGTGTGGTAGGGGAGATCGGAATTCCTGGTGTAgCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAA GGCGGATCTCTGGGCCATTACTGACgTcTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGgTAG TCCACGCCGTAAACGTTGGGAACTAGgTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCATTAAGTT CCCCGCCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCAGCGGAGCATG TGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCC CCTTGTGGTCGGTATACAGGTGGTGCATGGCTGTCGTCAGCTCGTGAGATGTTGGGTTAAGTCCCGCAAC GAGCGCAACCCTTGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGGGGTCAA CTCGGAGGAAGGTGGGGACGACGTCAAGTCATCCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCG GTACAATGAGCTGCGATGCCGCGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAAC TCGACCCCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTA AAGGTGGGACTGGCGATTG

16/30

[FIGURE 15 CONTINUED]

EN27

SEQ ID NO:12

TTAANACATGCAANTCGAACGATGAACCCNGTTTCGGTGGTGGATTAGTGGCGAACGGTGAGTAANANGTGGGCA ATTTGCCCTTCATTTTGGACAAGCCCTGGAAACGGGTTTAATACCGGATAACATTTTNTCCCGCATGGGANGGGG TTGAAAGNTCCGGCGGTGAAGGATGAGCCCGCGGCCTATNAGCTTGTTGGTGGGGTAATGGCCTACCCAAGGGAG ACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGAATGAGANACGGCCCAGAATCCTACGGGAGGCAGCA GTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGANGCCGCGTGAGGGATGACGGCCTTNGGGTTGTAA ACCTTTTTNAGCAGGGAAGAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATAAGTGCCAGCAGCC GCGGTAATAAGTAGGCCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTTGTAGGCGGCTTGTCANGTNGG ATGTGAAAGCCCGGGGNTTAACCCCGGGTTTGCATTTGATACGGGCTAGNTAGAGTGTGGTAGGGGAGATNGGAA TTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACT GACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGTTGGGAAC TAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCATTAAGTTCCCCGNCTGGGGAGTACGGCCGC AAGGCTAANACTCAAAGGAATTGACGGGGGCCCGNACAAGCAGCGGANCATGTGGCTTAATTCGACGCANCGCGA AGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCCTTGTGGTCGNTATACANGTGG TGCATGNCTGTCGTCACCTCGTGTCGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCNACCCTTGNTCTGTGTT GNCANCATGCCCTTCGGGGNTGATGGGGACTCACAGGANACTGNCCGGGGTCAACTCCGGANGAAGGTGGGTGAC GAAGTCAAGGTCATCATGNCCCCTTATGTCTTGGTGCTGCACACGTGC

EN28

SEQ ID NO:13

ACCGGAAAGCATCAGAGATGGTGCCCCCCTTGTGGTCGGTATACAGGTGGTGCATGGCTGTCGTCAGCTCGTGTC
GTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGGTTCTGTGTTTGGCCAGCATGCCCTTCGGGGTGAT
GGGGACTCACAGGAGACTGGCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAGTCATCATGCCCCTTATG
TCTTGGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGATGCCGCGAAGGCGGAGCGAATCTCAAAA
AAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGC
ATTGCTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACGTCACGAAAGTCGGTAACACCCGAAGC
CGGTGGTCCAACCCCTTGTGGGAGGGAGCTGTCGAAGGTGGGACTGGCGATTGG

17/30

[FIGURE 15 CONTINUED]

EN46

SEQ ID NO:16

ATGCAAGTCGAGCGGAAAGGCCCTTCGGGGTACTCGAGCGGCGAACGGGTGAGTAACACGTGAGTTAATCTGCCC CAGGCTCTGGATACCCACCGGAAAACGGTGATTAATACCGAATACGACAACCGATTTGCATGATCTGGTgGTGNA AAGTTTTTCGGCCTGGGATGTGCTTCGCGGCCTATCAGCTTGTTGGTGAGGTAATGGCTCACCCAAGGCTTCGAC GGTAGCCGGCCTGAGAGGGTGACCGNCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTG GGGAATATTGGACAATGGGCGGAAGCCTGATCCAGCAACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACC TCTTTCAGCACAGACGAAGCGCAAGTGACGGTATGTGCAGAAGAAGGACCGGCCAACTACGTGCCAGCAGCCGCG GTAATACGTAGGGTCCGAGCGTTGTCCGGAATTATTGGGCGTAAAGGGCTCGTAGGCGGTCTGTCGCGTCGGGAG TGAAAACCAGGTGCTTAACACCTGGCCTGCTTTCGATACGGGCAGNCTAGAGGTACNCAGGGGAGAATGGAATTC $\tt CGCTGAGGAGCGAAAGTGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACACCGTAAACGTTGGGCGCTA$ GGTGTGGGACACATTCCACGTGTTCCGTGCCGCAGCTAACGCATTAANCGCCCCGCCTGGGGAGTACGGCCGCAA NGCTAAAACTCANAGGAATTGACGGGGGCCCGCACAAGCGGGGAGCATGCGGATTAATTCGATGCAACGCGAAG ${\tt AACCTTACCTGGGTTTGACATACACCGGAAAGCCGTACAGATACGGCCCCTTTTAGTCgGTGTaCAGGTGGTGCA}$ TGGCTGTCgtCAgCtCgCTGTCGTGAGATGTtCGGGTTAAGTCCCGCAACGAGCGCAaCCCTCgTCCTATGTTGC ${\tt CagCAATTCGGTTGgGGACTCATAGGAgACTGCCgGGGGTCaACTCGGAGGAAGGTGGGGGATGACGTCAAGTCATC}$ ATGCCCCTTATGTCCAGGGCTTCACGCATGCTACAATGGCCGGTACAAAGGGCTGCGATCCCGTGAGGGTGAGCG AATCCCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTCGCTAGTAATCG CAGATCAGCAACGCTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACGTCACGAAAGTCGGCAAC ACCCGAAGCCANTGGCCCAACTCGTAAGAGAGGGAGCTGT

EN60

SEQ ID NO:18

ATGCAAGTNGAACGATGAANCCNTTTGGGGTGGATTAGTGGCGAACGGGTGAGTAANANGTGGGCAATTTGCCCT TCAATTTGGGACAAGCCCTGGAAACGGGGTNTAATACCGGATAACANTNTGTCCCGCATGGGACGGGGTTAAAAG CTCCGGCGGTGAAGGATGAGCCCGCGGCCTATNAGCTTGTTGGTGGGGTGATGGCCTACCAAGGCGACGACGGGT AGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGG AATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGCCTTCGGGTTGTAAACCTTT TTCAGCAGGGAAGAAGTGACGGTACCTGCAGAAGAGCGCCGGCTAAATANGTGCCAGCAGCCGGGTA ATANGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGTTTGTAGGCGGCTTGTCACGTNGGATGTGA GTGTAGCGGTGAAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGaTcTcTGGGCCATTACTGACGNT GAGGAGCGAAAGCGTGGGGAGCNAACAGNATTAGATACCCTGGTAGTCCAAGCCGTAAACGTTGGGAACTANGTG TTGGCGACATTCCACGTCGTCNNTGCCGCANCTAACGCATTAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCT AANACTCAAAGGAATTGANGNNGGCCCGCACAAGCAGCGGAGCATGTGGCTTANTTCNACGCANCGCGAAGAACC TTACCAAGGTTTGCCATATAcCGGAAAgCaTCAgAgATgGTGCCCCCCTtGTGGTCGGTATACAGqTGGTGCNTG GCTGTCGTCAgCTCGTGTCGTGACAtGTtGGTTAAgTCCCGTCAaCGAGqCGCAACCCTTGTTNTGTGTNGCCAG CATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGGGGTCAACTCGGAGGAAGGTGGGGACGACGTCAAG TCATCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGATGCCGCGAGGCG GAGCGAATCTCAAAAAGCCGGTNTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTCGGAGTTGCTAGT AATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACGTCACGAAAGTCG GTAACACCCGAAGCCGNTGG

[FIGURE 15 CONTINUED]

PM87

SEQ ID NO:24

GGCCCAGANATCCGNCTTCGCCACCGGTGTTCCTCCTGAATATCTGCGCATTTCACCGCTACACCAGGAATTCCG
ATCTCCCCTACCACACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACATCCGAC
GTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACCGCGGCTG
CTGGCACGTAATTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAGGGTTTACA
ACCCGAAGGCCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCCACTGCTG
NCTCCCGTANGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTCGNCCTCTCAGGCCGGCTACCGTCGC
CTTGGTAGGCCATTACCCCACCAACAAGCTGATANGCCGNGGGCTCATCCTTCANCGTCGGAGCTTTCAANCCCG
TCCATGCGGGACAGAGTGTTATCCGGTATTANACCCCGTNTCAGGGCTTGTCCANAGTGAAGGGCAGATNGCCAC
GTGTTATCACCGTTCGCCACTAATNACANCGAAACGGCTTATCGTNCGACTGCATGTGTTAACACNCGCAGCGTT
CGTCCTGAGCCAG

19/30

FIGURE 16

EN5

SEQ ID NO:3

GTAATGGCCCANAAAACCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTACACCAGGAAT TCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACTCGGGGTTAAGCCCCNAGCTTTCACATC CGACGTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGANAACGCTCGCACCCTACNTNTTACCGCG GCTGCTGGCNCGTNTTTAGCCGGTGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTNAAAAAAGGTT TACAACCCTANGGCCGT

EN6

SEQ ID NO: 4

EN7

SEQ ID NO:5

CCGCCTTCGCCACCGGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTACACCAGGAAATTCCNATCTCCCCTA
CCACACTCTANCTANCCCGTATCGAATGCAAACCCGGGGTTAANCCCCGGGCTTTCACACCCGACNTGACAAGCC
GCCTACAAACTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACNTATTACCGCGGCTGCTGGCACNTA
TTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAAAGGTTTACAACCCGAAGGC
CGTCATCCCTCACGCGGCGTCGCTCATCAGGCTTTCGCCCATTGTGCAATATTCCCCACTGCTGCCTCCCNTAG
GAATCTGGGCCGTGTCTCAATCCAGTGTGGCCGGTCCCCTCTCNGGCCGGCTACCGTCNTCCCTTGGTNACCATT
ANCTCACCAACAACTGATAGGNCGCGGGCTCATCTTCACGCGGGAACTTTCAACCACC

EN9

SEQ ID NO:6

GGCGGCGTGCTTAACACATGCAAGTCGAACGATGAAGCCCTTCGGGGTGGATTAGTGGCGAACGGGTGAGTAACA
CGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGTCTAATACCGGATACGATTCGGGAGGCAT
CTCCTGGtaCTGGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTTGTGGGTAATGGCCTACC
AAGGCGACGACGGGTAGCCGGCCTGAGAGGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGG
GAGGCAGCAGGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCACGCCCGCGTGAGGGATGACGCCTTC
GGGTTGTAAACCTCTTTCAGCAGGAAGAAGCGAAGAAGCGTACCTGCAGAAGAAGCGCCGGCTAACTACGTG
CCAGCAGCCGCGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAAGAGCTCGTAGGCGGCTTG
TCACGTCGGGTGTGAAAGCCCGGGGCTTAACCCCGGGTCTCGATCCGATACGGGCAGGCTAGAGTGTGGTAGGGG
AGATCGGAATTCCTGGTGTAACCCCGGGTCTCATCCGATACCGCTGGCGAAGCCGGATCTCTGG
GCCATTACTGACGCTGAGGAGCGAAATGCGCAGATATCAGGAGAATACCCTGGTAGTCCACGCCGTAAAC
GTTGGAACTAGGTGTTGGCGACATTCCACGTCGTCGTGCCCCAACGCATTAAGTTCCCCGCCTGGGGAGT
ACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCACAAGCAGCGGAGCATGTGGCTTAATTCGACG
CAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCGCCCACAAGCAGCATGTGGCTTAATTCGACG
CAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCGCCCAGAGATGGTGCCCCCCTTGTGGTCGGTAT
ACAGGTGGTGCATGGCTTGCCAGCTCGTTCGTGAGATGTTGGGTTAACCCCTTGT

[FIGURE 16 CONTINUED]

EN17

SEQ ID NO:8

CCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTACACCAGGAATTCC
NATCTCCCTACCACACTCTAGCTAGCCCGTATCAAATGCAAACCCGGGGTTAAGCCCCGGGCTTTC
ACATCCNACGTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCT
ACNTATTACCGCGGCTGCTGGCACNTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCT
NCTTCCCTGCTGAAANAGGTTTACAACCCAAAGGCCNTCATCCCTCNCCGGCNTCNTTGCNTCNGGC
TTNCNCCCATTGTTCAANNTTCCCCACTGCTNCCTCCCCTCGGAATCTGGGCCGNTGTCTCATTCCCN
TTNTGGCCGGTCCCCCTCNCAGGCCNGCTACCC

EN19

SEQ ID NO:9

CTCAGCGTCNGTAATGGCCCAAAAACCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTAC
ACCAGGAATTCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACCCGGGGTTAANCCCCGGGC
TTTCACATCCNACNTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGACAACGCTTGCNCCCTACTT
ATTACCGCGGCTGCTGGCACTTATTTAGCCGGCGCTTCTTCTCGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTN
AAAAAGGTTTACAACCCNAAGGCCGTCATCCCTCACGCGGCNTCGCTGCATCAGGCTTTCNCCCATTGTGCAATA
TTCCCCACTGCTGCCTCCCGTAGGATTCTGGGCCGTNTCTCATTCCCANTGTGGCCGGTCGCCCTCTCAGGCCGG
CTACCCGTCNTCNCCTTGGTAGGCCATTACCCCACCAACAAGCTNATAGGCCGCGGGCTCATCCTTCACCGCCGG
AAGCTTTCAACCCCNTCCATGCGGGANAAATTGTTNTCCGGTATTAAACCCCGTTTCCAGGGNTTGTCCCAAAAT
TGAAGGGGGGATTGNCCACTTTTTACTCACCCGTTCNCCNCTAATCCACCACC

EN26

SEQ ID NO:11

CCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTACACCAGGAATTCCNATCTCCCCTACC
GAACTCTANCCTGCCCGTATCNACTGCAAACCCGGGGTTAAGCCCCGGGCTTTCACAACCGACNTGACAAGCCGC
CTACAANCTCTTTACNCCCAATAATTCCGGACAACGCTTGCGCCCTACNTATTACCGCGGCTGCTGGCACNTATT
TAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAAAGGTTTACAACCCGAAGGCCG
TCNTCCCTCACGCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCCACTGCTGCCTCCCGTAGGA
TTCTGGGCCGTGTCTCANTCCCANTNTGGCCGGTCCCCTCTCAGGCCGGNTACCCGTCGTCCCTTGGTGAACCNC
TACCTCNCCAACAANCTGATAGGGCGCGGGCTCANCNTGCACGCCGGANCTTT

EN35

SEQ ID NO:14

21/30

[FIGURE 16 CONTINUED]

GGTGGTGCATGGCTGTCANCTCGTGTCGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTTCT
GTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGCCGGGGGTCAACTCGGAGGAAGGTGGGGA
CGACGTCAAGTCATCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGCTACAATGACCTGCGATG
CCGCGAGGCGGACCGAATCTCAACAAGCCCGTCTCATTCGGATTGCGGTCTGCAACtcCGACCCCATGAAGTCC
GACTtGCTAGTACTCGCACGTCAACATtGCTGCGCTGAATACGTCCCCGGGCCTTGTACACACCCCGTCACGT
CACGAAAGTCGGTAACACCCCGAAGCCGGTGGNCCAACCCCTTGTGGGAGGGAGCTGTCGAA

EN39

SEQ ID NO:15

ccgccttcgccaccggtgttcctcctgatatctgcgcatttcaccgctacaccaggaattccnatctcccctacc acactctagctancccgtatcnaatgcaaacccggggttaacccccgggctttcacacccnacntnacaanccgc ctacaaactctttacgcccaataattccggacaacgcttgcgccctacttattaccgcggctgctgcgcgcttatt tagccggcgcttcttctgcaggtaccgtcactttcgcttcttccctgctgaaaaaggtttacaacccgaaggcng tcatccctcacgcggcntcgctgcatcaggctttcgcccattgtgcaatattccccactgctgctcccgtagna ntctgggccgtntctcantcccagtgtggncggtcgccctctcaggccggctacccgtcgtcncctnggtnaacc attanntcaccaacaagctgataggccgcgggctcatccttcaccgcggagcttttaacccctgcccatgaaaa cagangtnttatccggtattanaacccgtttccaggg

EN57

SEQ ID NO:17

GTGCTTAACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACGGGTGAGTAACACGTG GGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTCTGTCCCGCATGGG ACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTTGGTGGGTAATGGCCTACCAAG GCGACGACGGGTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAG GCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCCTTCGGG TTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTACGTGCCA ${\tt GCAGCCGCGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGCTTGTCA}$ ${\tt TCGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATNCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGCC}.$ ATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGTT GGGAACTAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTGAACGCATTAAGTTCCCCGCCTGGGGAGTA: CGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCAGCGGAGCATGTGGCTTAATTCGACGC AACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCCTTGTGGTCGGTATA CAGGTGGTGCATGGCTGTCAGCTCGTGTGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCTTGTT $\tt CTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACTGcCGGGGTCAACTCGGAGGAAGGTGGG$ GACGACGTCAAGTCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCTGCGA TGCCgCGAgGCGAgCgAATCTCAAAAAGCCGGTCTCAqTTCGGATTGGGGTCTGCAACtCGACCCCATGAAGTC GGAgTTGCTAgTAATCgCAgATCAgCATTGCTGcGGTGAATACGTTCCCGGGCCTTGTACACACCGCCGTCAcGT CACGAAAGTCGGTaACACCCGAAGCCGGTGGCCCAACCGCCTTGTGGGAAGGGAACTTTCCA

22/30

FIGURE 17

SE1

SEQ ID NO:19

GAACGATGAAGCCGTTTCGGTGGTGGATTAGTGGCGAACGGTGAGTAAAAGTGGCAATTTNCCCTTCATTTTGGA CAAGCCCTGGAAACGGGTTTAANACCGGATAACATTNTGTCCCGCATGGGACGGGGTTGAAAGNTCCCGGCGGTG AAGGATGAGCCCGCGGCNTATCAGCTTGTTGGTGGGGTAATGGCCTACCAAGGCGACGACGGGTAGCCGGCCTGA GAGGGCGACCGGCCACACTGGGANTGAGACACGGCCCAGACTCCTACGGGAGGCAGCAGTGGGGAATATTGCACA ATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACCTNTTTCAGCAGGGA AGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAAATANGTGCCAGCAGCCGCGGTAATANGTAGGGC GCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTTGTAGGCGGCTTGTCANGTCGGATGTGAAAGCCCGGGGC TTAACCCCGGGTTTGCATTCGATACGGGCTAGTTAGAGTGTGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTG AAATGCGCAGATATCAGGAGGAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAA GCGTGGGGAGCNAACAGGATTAGATACCCTGGTAGTCCACGCCGTAAACGTTGGGAACTAGGTGTTGGCGACATT CCACGTCGTCGCTGCCGCCTAACGCATTAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAG GAATTGACGGGGGCCCGCACAAGCAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAACAACCTTACCAAGGCT ${\tt TGACATATACCGGAAAGCATCANAGATGGTGCCCCCCTTGTGGTCGTATACANGTGGTGCATGGCTGTCGTCAG}$ $\tt CTCGTGTCGTGAGATGTTGGGTTANGTCCCGCAACGAGCGCNACCCTTGTTCTGTTGTGTCGNCNAGCATGCCCTTCG$ ${\tt NGGTGATGGGGACTCACANGAGACTGNCGGGGTCCACTCGGAGGAAGGTGGCGACNACGTCANNTCATCATGCCC}$ CCTTATGTCTTGGGNCTGGCCACGTGCNACNATGGCC

SE2

SEQ ID NO:20

GCTGGCGGCGTGCTTAACACATGCAAGTCGAACGATGAAGCCGCTTCGGTGGTGGATTAGTGGCGAACGGGTGAG TAACACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTCTGTC CCGCATGGGACGGGGTTGAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTTGGTGGGGTAATG GCCTACCAAGGCGACGACGGCTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACT CCTACGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGAC GGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAA CTACGTGCCAGCAGCCGCGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGG TCTCTGGGcCAtTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCAAGC CGTAAACGTTGGGAACTANGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCATTAAGTTCCCCGTC CTGGGGAGTACGGCCGCNAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCAGCGGAGCATGTGGCTT ANTTCGACGCNACGCGAAGAACCTTNCCAAGGCTGACATATACCGGAAAGCATCAcAGATGGTGCCCCCCTTGTG ${\tt GTCGGTATACAGGgTGGTGCATGGCTGTtCGtCaGCTCGTGTCgtGAGATGTTGGGTTAaGTCCCGCAAAGAGCG}$ CAACCGTGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGACTCAcAcGAGACTGTCNGGGTCAACTCgga ${\tt GGAAgGTGGgGACGACgTCAAGTtCATCCCCCTTATGTCTTGGGCTGCACACGNGCTACAATGGCCGGTACA}$ ATGAGNNGGGATGCCGCGAGGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTGACC CCATGAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTNCCCGGGCCTNGTACACACC ACCCGTCACGTCACGAAAGTCGGTAACACCCTAAGCCGGTGNCCCAACCCCTTNTGGGAGG

FIGURE 18

PM36

SEQ ID NO:21

CCAGANATCCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTACACCAGGAATTCCGATCT
CCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACATCCGACGTGA
CAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACCGCGGCTGCTGG
CACGTAGTTAGCCGGCGCTTCTTCTCCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAGGTTTACAACCC
GAAGGNCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCCACTGCTGCCTC
CCGTAGGAGTCTGGGNCGTGTTCAATNCCAGTGGTGGGCCGGTCGCCCTCTCAGGNCGGCTACCGTCGTCGCCTT
GGTAGGCATTACCACAACAAGCTGATAGGCGGGGGTCATCCTTCAACGCCGGAGCTTCAAACCCGTCCATGCGGG
ACAAGTGTATCCGGTATTAAACCC

PM40

SEQ ID NO:22

TCAGTNATGGCCCAGAANGATCCGNCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTACACCA
GGAATTCCGATCTCCCCTACCACACTCTAACTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTC
ACATCCGACGTGACAAGCCGCCTACGAGCTCTTNACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTA
CCGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTNACTTTCGCTTCTTCTCCTGCTGAAAG
AGGTTTACAACCCGAAGGCCGTCNTCCCTCACGCGGCGTCGCATCAGGCTTTCGCCCATNGTGCANTATTCC
CCACTGNTGNCTCCCGTANGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTCGNCCTCTCAGGCCGGCTAC
CGTCGTCGCCTTGGTAGGNCATTACCCACCAACAAGCTGATANGTCGNGGGCTCATCCTTCACCGNCGGAGNTTT
AACCCCGTNCATGCGGGACAGAGTGTTATCCGGTATTANACCCGTATNCAGGGCTTGTCCCATAGTGAAGGGNAG
ATNGCCACGTGTTATCACCGTTCGNCACTAATNATCANCGAANCGGCTTCATCGTTCGACTTGCATGTGTTA

PM41

SEQ ID NO:23

PM171

SEQ ID NO:25

[FIGURE 18 CONTINUED]

PM185

SEQ ID NO:26

TCAGTAATGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCCTCCTGGATATCTGCGCATTTCACCGCTACACCAG
GAATTCCGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCA
CATCCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTAC
CGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGA
GGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCC
CACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCAGTGTGGCCGGTCGCCTCTCAGGCCGGCTACC
CGTCGTCGCCTTGGTAGGCCATTACCCCACCAACAAGCTGATAGGCCGCGGGCTCATCCTTCACCGNCGGAGCTT
TAACCCCGTCCCATGCGGGACAGAGTGTTATCCGGTATTAGAACCCGTTTCCAGGGCTTTGTCCCAGAGTGAAGGG
CAGATTGCCACGTGTTACTCANCCGTTCGNCACTAATCANCAACGAAGCGGCTTCATCGTTCGACTTGCATGTGT
TAAGCACGCCGNCAGCGTTCGTCCTGAGCCAGGATC

PM208

SEQ ID NO:27

TCAGTATCNGCCCAGAGATCCGCCTTCGCCACCGGTGTTTCCTCCTGATATCTGCGCATTTCACCGCTACACCAG
GAATTCCGATCTCCCCTACCGAACTCTAGCCTGCCCGTATCGACTGCAGACCCGGGGTTAAGCCCCGGGCTTTCA
CAACCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTAC
CGCGGCTGCTGGCACGTAGTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGA
GGTTTACAAACCGAAGGCCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCC
CACTGGTGNCTCCCGTANGAGTCTGGGGCGGTGTCTCANTCCAGTGTGGGCGGTCGCTCTCAGGGCGCTACCGT
CGTCGCTTGGTGAGNCACTACTCACAACAAGCTGATAGGCGCGGGCTCATCTGGAACGGCGGAGCTTTACAC

PM228

SEQ ID NO:28

TCAGTAATGGCCCAGANATCCGNCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTACACCAGG AATTCCGATCTCCCCTACCACACCTCTAACTAGCCCGGTGTTCCCCTGATATCTGCGCCTTTAAGCCCCGGGCTTTCAC AATTCCGACGTGACAAGCCGCCTACCACCTTTACGCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACC GCGGCTGCTGGCACGTAGTTAGCCGGGCGTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAG GTTTACAACCCGAAGGCCGTCATCCCTCACGCGGGCGTCGCTGCATCAGGCCTTTCGCCCATTGTGCAATATTCCCC ACTGCTGCCTCCCGTANGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCCGGTCGCCCTCTCAGGCCGGCTACCG TCGTCGCCTTGGTAGGCCATTACCCACCAACAAGCTGATANGNCGNGGGCTCATCCTTCACCGNCGGAGCTTTCAANCCCGTCCCATGCGGGACAGAGTGTTATCCGGTATTAAACCCGTNTCCAGGGCTTGTCCATAGTGAAGGGCAGA TTGCCAAGTGTTATCANCCACTAATCATCANCGAAGCGGCTTCATCGTTCGACTGCATGTGTT

PM252

SEQ ID NO:29

TCCTCAGNATCAGTAATGGCCCAGAGATCCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGC
TACACCAGGAATTCCGATCTCCCCTACCACACTCTANCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCG
GGCTTTCACATCCGANGTGACAAGCCGCCTACGAGCTCTTTACGCCCAATAATTCCGGACAANGCTTGCGCCCTA
CGTATTACCGCGGNTGCTGGCACGTAGTTAGCCGGCGCTCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTG
CTGAAAGAGGTTTACAACCCGAAGGCCGTCATCCCTCACNCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCA
ATATTCCCCACTGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAATCCCANTGTGGCCGGTCGCCCTCTCANGC
CGGCTACCGTCGTCGCTTGGTAGGCCATTACCCCACCAACAAGCTGGATAGGNCGGGGGCTCATTCTTCACCGCC
GGAAGCTTTAANCCCGTCCATGCGGGANANAGTGNATCCCNGTATTAAACCCNGTTTCAGGGCTTGTCCANAGTG
AAGGGNGATTGCCCNAGTGTTTATCNCCCGTTCGCCANTAATCNACAACGAAAGCGGNTTCNTCGNTTCGACTTG

25/30

[FIGURE 18 CONTINUED]

PM342

SEQ ID NO:30

TAATGGCCCAGAANATCCGCCTTCGCCACCGGTGTTCCTCCTGAATATCTGCGCATTTCACCGCTACACCAGGAA
TTCCGATCTCCCCTACCACACTCTAGCTAGCCCGTATCGAATGCAGACCCGGGGTTAAGCCCCGGGCTTTCACAT
CCGACGTGACAAGCCGCCTACGAGCTCTTTACGCCCCAATAATTCCGGACAACGCTTGCGCCCTACGTATTACCGC
GGCTGCTGGCACGTAGTTAGCCGGCGTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGCTGAAAGAGGT
TTACAACCCGAAGGCCGTCATCCCTCACGCGGCGTCGCTGCATCAGGCTTTCGCCCATTGTGCAATATTCCCCAC
TGCTGCCTCCCGTAGGAGTCTGGGCCGTGTCTCAGTCCCAGTGTGGCGGTCGCCCTCTCAGGCCGGNTANCCGTC
GTCGCCTTGGTANGCCATTANCCCACCAACAAGCTGATANGCCGNGGGCTCATCCTTCANCGCCGGAGCTTTTAA
CCCCGTCCCATGCGGGACAGAGTGTTATCCGGTATTAGATCCCGTNTCCAGGGCTTGTNCATAGTGAAGGGCANA
TTGCCACGTGTTACTCANCCGTTCGC

FIGURE 19

ENZ

GGCGGCGTGCTTAACACATGCAAGTCGAACGATGAACCACTTCGGTGGGGATTAGTGGCGAACGGGTGAGTAACA CGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTtAATACCGGATAACACTCCCACTCTC $\tt CTGAGTGGGGGTTAAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTTGGTGAGGTAATGGCTC$ ACCAAGGCGACGACGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTA TTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTAC GTGCCAGCAGCCGCGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTAGGCGGC GGGAGATCGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATTCAGGAGGAACACCGGTGGCGAAGGCGGATCT CTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGT AAACGGTGGGAACTAGGTtGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCATTAAGTTCCCCGCCTG GGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCAGCGGAGCATGTGGCTTAAT TCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACACCGGAAAGCATCAGAGATGGTGCCCCCCTTGTGGT CCTTGTTCTGTGTGCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGAGACCGCCGGGGTCAACTCGGAGGAA GGTGGGGACGTCAAGTCATCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCAGGTACAATGAG $\tt CTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCTGTCTCAgTTCGGATTGGGGTCTGCAACTCGACCCcaT$ GAAGTCGGAGTTGCTAGTAATCGCAGATCAGCATTGCTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCC CTGGCGATTGGG

EN10

GAGTTTGATCNNGGCTCAGACGAACGCTGGCGGCGTGTTAACACAANCCAAGTCGAANGNTGAACCACTTCGTTG ${\tt GGATTAGTGCGAACGGTGNTAACACGNTGGCAATGTGCCCTTCACTNTGGGACAAGNCCTGGAAACCGGGTTCTA}$ ATACCGGATACCACTACCCGCAGGCATCTGTGGTGTTTGAAAGCTCCGCCGTTGAAGGATGAGCCCGCGGCCTAT CAGCTTGTTGGTGAGGTAATGGCTCACCCAAGGCGACGACGGATAGCCGGCCTGAGAGGGGGGCCACACTG GGACTGAGACACGGCCCAGACTCCTACGGAGGCAGCAGTGGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAG CGACGCCGCGTGAGGGATGACGGCCTTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACC TGCAGAAGAAGCGCCGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTAT TGGGCGTAAAGAGCTCGTAGGCGGCTTGTCACGTCGGGTGTGAAAGCCCGGGGCTTAACCCCGGGTCTGCATTCG ATACGGGCTAGCTAGAGTGTGGTAGGGGAGATCGGAATTCCTGGTGTACCCGGTGAAATGCGCAGATATCAGGAG GAACACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGAT TAGATACCCTGGTAGTCCACGCCGTAAACGGTGGGAACTAGGTGTTGGCGACATTCCACGTCGTCGTGCCGCAG $\tt CTAACGCATTAAGTTCCCCGCCTGGGGAGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCAC$ AAGCAGCGGAGCATGTGGCTTAATTCGACGCAACGCGAAGAACCTTACCAAGGCTTGACATACGCCGGAAAGCAT CAGAGATGGTGCCCCCCTTGTGGTCGGTGTACAGGTGGTGCATGGCTGTCGTCGTGAGATGTTGG GTTAAGTCCCGCAACGAGCGCAAcCCTTGTCCTGTGTTGcCAGCATGCCCTTCGGGGTGATGGGGACTCACAGGA GACCGCCGGGGTCAACTCGGAGGAAGGTGGNGACGACGTCGAGTCATCATGCCCCTTATGTCTTGGGGCTGCACA CGTGCTACNATGggCaGGTACAATGAGCTGCGATACCGTGAGGTGGAGCgCATCTnnnnnAGCctGTCTCAGTTC GgATTGGGGTCTGcAACTCGACCCCaTGAAGTCGGAGTTGCTAgATAATCgCAGATCAGCATTGCtGCGqTGAAT ACGTtCCCGGGCCTTGTACACCCCCCCCCTCACGTCACGAAAGTCGGTAACACCCGAAGCCGGTGGCCCAACCCC TTGTGGGAGGGAGCTGTCGAANGTGG

27/30

[FIGURE 19 CONTINUED]

EN22

TCCTCAGCGTCAGTAATGGCCCAAAAACCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCT
ACACCAGGAATTCCNATCTCCCCTACCACACTCTAGCTAGCCCGTATCNAATGCAAACCCGGGGTTAANCCCCGG
GCTTTCACATCCNACNTGACAAGCCGCCTACAANCTCTTTACGCCCAATAATTCCGGACAACNCTTGCGCCCTAC
TTATTACCGCGGCTGCTGGCACTTATTTAGCCGGCGCTTCTTCTGCAGGTACCGTCACTTTCGCTTCTTCCCTGC
TGAAAAAGGTTTACAACCCNAAGGCCGTCATCCCTCACGCGGCNTCNCTGCATCAGGCTGTCCCCATTGTGCAA
TATTCCCCACTGCTGCCTCCCGTAGGATTCTGGGCCGTNTCTCANTCCCANTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTCGTCNCCTTGGTAGGCCATTACCCCNCCAACAANCTGATAGGCCGCGGGGCTCNTCCTTCACCGCC
GGAGCTTTCAACCCCGTCCCATGCGGGANAAANTGTTNTCCGGTATTAAAACCCGTTTCCAGGGNTTGTCCAAAA
TTGAAGGGNANATTGCCCACTTTTNNTCACCCGTTCCCCACCACCACA

EN30

 $\tt TGGNGGNGTGCTTAACACATGCAAGTCGAACGATGAANCCTTTCGGGGTGGATTAGTGGCGAACGGGTGAGTAAC$ ACGTGGGCAATCTGCCCTTCACTCTGGGACAAGCCCTGGAAACGGGGTCTAATACCGGATAACACTCTGTCCCGC ${\tt ATGGGACGGGGTTAAAAGCTCCGGCGGTGAAGGATGAGCCCGCGGCCTATCAGCTTGTTGGTGGGGTGATGGCCT}$ ACCAAGGCGACGACGGCTAGCCGGCCTGAGAGGGCGACCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTA CGGGAGGCAGCAGTGGGGAATATTGCACAATGGGCGAAAGCCTGATGCAGCGACGCCGCGTGAGGGATGACGGCC TTCGGGTTGTAAACCTCTTTCAGCAGGGAAGAAGCGAAAGTGACGGTACCTGCAGAAGAAGCGCCGGCTAACTAC GTGCCAGCAGCCGCGTAATACGTAGGGCGCAAGCGTTGTCCGGAATTATTGGGCGTAAAGAGCTCGTANGCGGC GGAGATCGGAATTCCTGGTGTANCGGTGAAATGCGCAGATATCAAGAGGAACANCGGTGGCGAANGCGGATCTCT GGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAGATACCCTGGTAGTCCACGCCGTAA ACGTTGGGAACTAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTAACGCATTAAGTTCCCCGCCTGGGG AGTACGGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCAGCGGAGCATGTGGCTTAATTCG ACGCAACGCGAAGAACCTTACCAAGGCTTGACATATACCGGAAAGCATCAGAGATGGTGCCCCCCTTGTGGTCGG ${\tt TATACAGGTGGTGCATGGCTCGTCGTCGTGTCGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCAACCCT}$ TGTTCTGTGTTGCCAGCATGCCCTTCGGGGTGATGGGGGACTCACAGGAGACTGCCGGGGTCAACTCGGAGGAAGG TGGGGACGACGTCAAGTCATCATGCCCCTTATGTCTTGGGCTGCACACGTGCTACAATGGCCGGTACAATGAGCT GCGATGCCGCGAGGCGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGcAACTCGACCCcATGA ${\tt AGTCGgAGTTGCTAgTAATCgCAgATCAgCATTGCTGCGGTGAATACgTTNCCGGGCCTtGTACAcACCGCCCGT}$ CTGGCGATTGG

EN43

CTCAGCGTCACTATCGGCCCAAAANACCGCCTTCGCCACCGGTGTTCCTCCTGATATCTGCGCATTTCACCGCTA
CACCAGGAAATTCCANTCTCCCCTACCGAACTCTANCCTGCCGTATCAACCGCAGGCTTGGGGTTAAGCCCCAA
TTTTTCACGGTCAACGCNACAAGCCGCCTACAAGCTCTTTACGCCCAATAAATCCGGACAACGCTCGCACCCTAC
TTCTTACCGCGGCTGCTGGCACTTATTTGGCCGGTGCTTCTTCTGCAGGTACCGTCACTCTCGCTTCGTCCTGC
TNAAAAAGGTTTACAACCCGAAGGCCGTCATCCCTCACGCGGCNTCGCTGCATCAGGCTTCCGCCCATTGTGCAA
TATTCCCCACTGCTGCCTCCCGTAGGATTCTGGGCCGTNTCTCANTCCCAGTGTGGCCGGTCGCCCTCTCAGGCC
GGCTACCCGTCGTCGCCTTGGTAGGCCATCACCCCACCAACAAGCTGATAGGCCGCNAAGCCCATCCCAAGCCGA
AAAACTTTCCACCACCAGCCATGCGGCCAAAAATTCCTATTCGGTATTAGCCCCCGTTTCCNAAGGTTNTCCCAAA
GCTTGGGGCAGGTTGCTCACTTTTTACTCACCCGTTCCCGCTCAATTACCCCNAAGGGGNTTTCCCTCAACTTGC
AT

28/30

[FIGURE 19 CONTINUED]

EN47

AACACGTGAGTAATCTGCCCCAGGCTCTGGGATAGCCACCGGAAACGGTGATTAATACCGGATACGACAACCGAT TGCATGATCTGGTTGTGGAAAGTTTTTCGGCCTGGGATGTGCTCGCGGCCTATCAGCTTGTTGGTGAGGTAATGG $\tt CTCACCAAGGCTTCGACGGCTAGCCGGCCTGAGAGGGTGACCGGCCACACTGGGACTGAGACACGGCCCAGACTC$ CTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGGCGGAAGCCTGATCCAGCAACGCCGCGTGAGGGATGACG GCCTTCGGGTTGTAAACCTCTTTCAGCACAGACGAAGCGCAAGTGACGGTATGTGCAGAAGAAGGACCGGCCAAC TACGTGCCAGCAGCCGCGGTAATACGTAGGGTCCGAGCGTTGTCCGGAATTATTGGGCGTAAAGGGCTCGTAGGC ${\tt GGTCTGTCGCGTCGGGAGTGAAAACCAGGTGCTTAACACCTGGCCTGCTTTCGATACGGGCAGACTAGAGGTACT}$ ${ t TCTCTGGGAGTATCCTGACGc{ t TTGAAGAGCGAAAGTGTGGGGAGCGAACAGGATTAGATANTNTGGTAGTCCACA}$ ${\tt CCGTAAACgTTGGGCGcTAGgTGTGGGACACATTCCACGTGTTCCGTGCCGCAGCTAACGCATTAAGCGCCCCGC}$ CTGGGGAGŤACGCCGCAAGGCTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCGGCGGAGCATGCGGATT AATTCGATGCAACGCGAAGAACCTTACCTGGGTTTGACATACACCGGAAAGCCGTAGAGATACGGCCCCTTTTAG GATGACGTCAAGTCATGCCCCTTATGTCCAGGGCTTCACGCATGCTACAATGGCCGGTACAAAGGGCTGCGA TCCCGTGAGGGTGAGCGAATCCCAAAAAGCCGGTCTCAGTTCGGATTGGGGTCTGCAACTCGACCCCATGAAGTC GGAGTCGCTAGTAATCGCAGATCAGCAACGCTGCGGTGAATACGTTCCCGGGCCTTGTACACACCGCCCGTCACG TG

EN59

GGGNATTAGTGGGGAACGGGTGAGTAAAANGTGGCCANTTTCCCCTGNATTTTGGACANCCCCNGGAAANGGNTT NTAAAACNGGATANTGACCACCTTGGCATCCAAGTTTTNGAAACTTCCGGCGGTGCAGGATGAGCCNGCGGCNTA TNAGCTTGTTGGNGAGGTAATGGNTCACCAAGGGANGACGGGTAGCCGGCCTGAGAGGGGACCNGCCACANTGGG ${\tt ANNCCGCGTGAGGGANGACGGCCTTNGGGTTGTAAACNTTTTTNAGCAGGGAAGAGCGAAAGTGACGGTACCTG}$ CAGAAGAAGCGCCGGCTAAATAAGTGCCAGCAGCCGCGGTAATAAGTAGGGNGCGAGCGTTGTNCGGAATTATTG ${\tt GGNGTAAAGAGTTTGTAGGCGGNTTGTNAAGTNGGTTGTGAAAGCCCGGGGNTTAACCCCGGGTTTGCAGTTGAT}$ ACGGGCAGGNTAGAGTTCGGTAGGGGAGATNGGAATTCCTGGTGTAGCGGTGAAATGCGCAGATATCAGGAGGAA CACCGGTGGCGAAGGCGGATCTCTGGGCCATTACTGACGCTGAGGAGCGAAAGCGTGGGGAGCGAACAGGATTAG ATACCCTGGTAGTCCACGCCGTAAACGGTGGGAACTAGGTGTTGGCGACATTCCACGTCGTCGGTGCCGCAGCTA CTACAATGGCCGGTACAAAGAGCTGCGATACCGTGAGGTGGAGCGAATCTCAAAAAGCCGGTCTCAGTTCGGATT GGGGTCTGCAACTCGANCCCATGAANTCGGAGTTGCTAATTAATCGCAAAATCAAGCATTGCTGGCGGTGAATAC GTTCCC

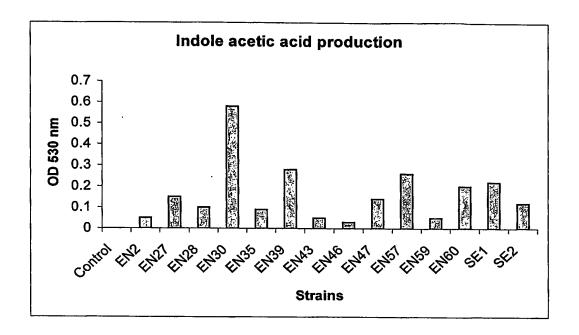
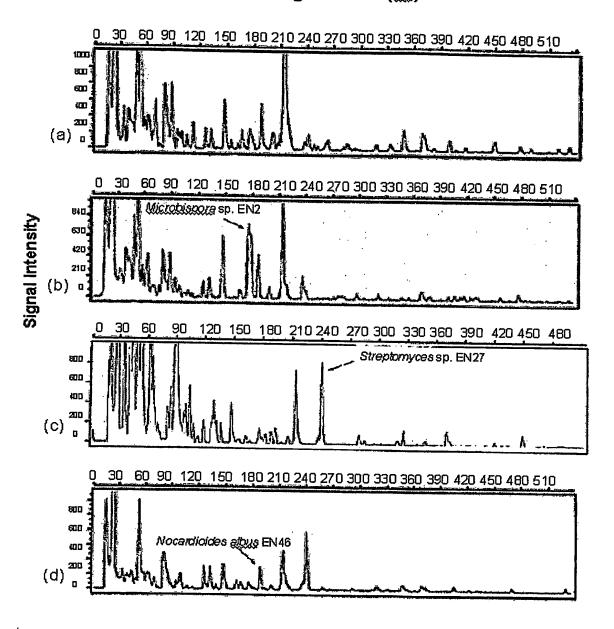


Figure 20

Figure 21





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